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## WHAT IS CLAIMED IS:

- 1. An isolated nucleic acid molecule consisting of a polynucleotide having a nucleotide sequence selected from the group consisting of:
  - a) a polynucleotide fragment of SEQ ID NO:1 or a polynucleotide fragment of the cDNA sequence included in ATCC Deposit No:PITA-2682, which is hybridizable to SEQ ID NO:1;
  - b) a polynucleotide encoding a polypeptide fragment of SEQ ID NO:2 or a polypeptide fragment encoded by the cDNA sequence included in ATCC Deposit No:PTA-2682, which is hybridizable to SEQ ID NO:1;
  - a polynucleotide encoding a polypeptide domain of SEQ ID NO:2 or a polypeptide domain encoded by the cDNA sequence included in ATCC Deposit No:PTA-2682, which is hybridizable to SEQ ID NO:1;
  - a polynuc eotide encoding a polypeptide epitope of SEQ ID NO:2 or a polypeptide epitope encoded by the cDNA sequence included in ATCC Deposit No:PTA-2682, which is hybridizable to SEQ ID NO:1;
  - e) a polynucleotide encoding a polypeptide of SEQ ID NO:2 or the cDNA sequence included in ATCC Deposit No:PTA-2682, which is hybridizable to SEQ ID NO:1, having biological activity;
  - f) a polynucledtide which is a variant of SEQ ID NO:1;
  - g) a polynucleo ide which is an allelic variant of SEQ ID NO:1;
  - h) a polynucleotide which encodes a species homologue of the SEQ ID NO:2;
  - i) a polynucleotide which represents the complimentary sequence (antisense) of SEQ ID NO:1;

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- j) a polynucleotide corresponding to nucleotides 4 to 954 of SEQ ID NO:1;
- k) a polynucleotide corresponding to nucleotides 1 to 954 of SEQ ID NO:1; or
- a polynucleotide capable of hybridizing under stringent conditions to any one of the polynucleotides specified in (a)-(k), wherein said polynucleotide does not hybridize under stringent conditions to a nucleic acid molecule having a nucleotide sequence of only A residues or of only T residues.
- 2. The isolated nucleic acid molecule of claim 1, wherein the polynucleotide fragment comprises a nucleotide sequence encoding a G-protein coupled receptor protein.
- 3. The isolated nucleic acid molecule of claim 1, wherein the polynucleotide fragment comprises a nucleotide sequence encoding the sequence identified as SEQ ID NO:2 or the polypeptide encoded by the cDNA sequence included in ATCC Deposit No:PTA-2682, which is hybridizable to SEQ ID NO:1.
- 4. The isolated nucleic acid molecule of claim 1, wherein the polynucleotide fragment comprises the entire nucleotide sequence of SEQ ID NO:1 or the cDNA sequence included in ATCC Deposit No:PTA-2682, which is hybridizable to SEQ ID NO:1.
- 5. The isolated nucleic acid molecule of claim 2, wherein the nucleotide sequence comprises sequential nucleotide deletions from either the C-terminus or the N-terminus.
- 6. The isolated nucleic acid molecule of claim 3, wherein the nucleotide sequence comprises sequential nucleotide deletions from either the C-terminus or the N-terminus.
- 7. A recombinant vector comprising the isolated nucleic acid molecule of claim 1.

- 8. A method of making a recombinant host cell comprising the isolated nucleic acid molecule of claim 1.
  - 9. A recombinant host cell produced by the method of claim 8.
  - 10. The recombinant host cell of claim 9 comprising vector sequences.
- An isolated polypeptide comprising an amino acid sequence at least 95% identical to a sequence selected from the group consisting of:
  - a) a polypeptide fragment of SEQ ID NO:2 or the encoded sequence included in ATCC Deposit No:PTA-2682;
  - b) a polypeptide fragment of SEQ ID NO:2 or the encoded sequence included in ATCC Deposit No:PTA-2682, having biological activity;
  - a polypeptide domain of SEQ ID NO:2 or the encoded sequence included in ATCC Deposit No:PTA-2682;
  - d) a polypeptide epitope of SEQ ID NO:2 or the encoded sequence included in ATCC Deposit No:PTA-2682;
  - e) a full length protein of SEQ ID NO:2 or the encoded sequence included in ATCC Deposit No:PTA-2682;
  - f) a variant of SEQ ID NO:2;
  - g) an allelic variant of SEQ ID NO:2;
  - h) a species homologue of SEQ ID NO:2; or
  - i) a polypeptide corresponding to amino acids 2 to 318 of SEQ ID NO:2.
- 12. The isolated polypeptide of claim 11, wherein the full length protein comprises sequential amino acid deletions from either the C-terminus or the N-terminus.
- 13. An isolated antibody that binds specifically to the isolated polypeptide of claim 11.
- 14. A recombinant host cell that expresses the isolated polypeptide of claim
  11.

- 15. A method of making an isolated polypeptide comprising:
  - a) culturing the recombinant host cell of claim 14 under conditions such that said polypeptide is expressed; and
  - b) recovering said polypeptide.
- 16. A polypeptide produced by claim 15.
- 17. A method for preventing, treating, or ameliorating a medical condition, comprising administering to a mammalian subject a therapeutically effective amount of the polypeptide of claim 11 or the polynucleotide of claim 1.
- 18. A method of diagnosing a pathological condition or a susceptibility to a pathological condition in a subject comprising:
  - a) determining the presence or absence of a mutation in the polynucleotide of claim 1; and
  - b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or absence of said mutation.
- 19. A method of diagnosing a pathological condition or a susceptibility to a pathological condition in a subject comprising:
  - a) determining the presence or amount of expression of the polypeptide of claim 11 in a biological sample; and
  - b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or amount of expression of the polypeptide.
  - 20. A gene corresponding to the CDNA sequence of SEQ ID NO:2.
- 21. A method of identifying an activity in a biological assay, wherein the method comprises:
  - a) expressing the HGPRBMY4 sequence as set forth in SEQ ID NO:2 in a host cell having; and
  - b) measuring the resulting activity of the expressed HGPRBMY4.

- 22. A method for identifying a binding partner to the polypeptide of claim 11 comprising:
  - a) contacting the polypeptide of claim 11 with a binding partner; and
  - b) determining whether the binding partner effects an activity of the polypeptide.
- 23. A method of identifying a compound that modulates the biological activity of HGPRBMY4, or a GPCR, comprising:
  - a) combining a candidate modulator compound with a host cell containing a vector according to claim 7, wherein HGPRBMY4 is expressed by the cell; and
  - b) measuring an effect of the candidate modulator compound on the activity of the expressed HGPRBMY4.
- 24. A compound that modulates the biological activity of human HGPRBMY4 as identified by the method according to claim 21, 22, or 23.
  - 25. The method of claim 22 wherein said binding partner is a peptide.
- 26. A method of treating a disease, disorder, or condition related to the colon, breast, ovaries, or immune system, comprising administering the G-protein coupled receptor polypeptide or homologue according to claim 11 in an amount effective to treat the lung-, colon-, brain-, heart-, or prostate-related disorder.
- 27. The polynucled tide of claim 2, further comprising a polynucleotide localized in lung, colon, brain, prostate, heart, colon carcinoma, or lung carcinoma cell lines.
- 28. The polypeptide of claim 11, further comprising a polypeptide expressed in lung, colon, brain, prostate, heart, colon carcinoma, or lung carcinoma cell lines.
  - 29. A cell comprising NFAT/CRE and the polypeptide of claim 11.
  - 30. A cell comprising NFAT G alpha 15 and the polypeptide of claim 11.

- 31. A method of screening for candidate compounds capable of modulating activity of a G-protein coupled receptor-encoding polypeptide, comprising:
  - a) contacting a test compound with the cell of claim 29 or 30; and
  - b) selecting as candidate modulating compounds those test compounds that modulate activity of the G-protein coupled receptor polypeptide.
- 32. The method according to claim 31, wherein the candidate compounds are agonists or antagonists of G-protein coupled receptor activity.
- 33. The method according to claim 32, wherein the candidate compounds are peptides.
- 34. The method according to claim 32, wherein the polypeptide activity is associated with the lung, colon, brain, heart, or prostate.

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